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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/459,062

DATE: 06/19/2001

TIME: 12:40:34

Input Set : A:\17634341.app

Output Set: N:\CRF3\06192001\I459062.raw

ENTERED

3 <110> APPLICANT: Tao, Tao
 4 Skiadopoulos, Mario H.
 5 Collins, Peter L.
 6 Murphy, Brian R.
 8 <120> TITLE OF INVENTION: CONSTRUCTION AND USE OF RECOMBINANT PARAINFLUENZA
 9 VIRUSES EXPRESSING A CHIMERIC GLYCOPROTEIN
 11 <130> FILE REFERENCE: 17634-000340US
 13 <140> CURRENT APPLICATION NUMBER: 09/459,062
 C--> 14 <141> CURRENT FILING DATE: 2001-06-04
 16 <150> PRIOR APPLICATION NUMBER: 09/083,793
 17 <151> PRIOR FILING DATE: 1998-05-22
 19 <150> PRIOR APPLICATION NUMBER: 60/059,385
 20 <151> PRIOR FILING DATE: 1997-09-19
 22 <150> PRIOR APPLICATION NUMBER: 60/047,575
 23 <151> PRIOR FILING DATE: 1997-05-23
 25 <160> NUMBER OF SEQ ID NOS: 57
 27 <170> SOFTWARE: PatentIn Ver. 2.1
 29 <210> SEQ ID NO: 1
 30 <211> LENGTH: 42
 31 <212> TYPE: DNA
 32 <213> ORGANISM: Artificial Sequence
 34 <220> FEATURE:
 35 <223> OTHER INFORMATION: Description of Artificial Sequence: Flanking
 36 sequence of measles HA gene insert for N-P and P-M
 37 junctions.
 39 <400> SEQUENCE: 1
 40 cttaagaata tacaaataag aaaaacttag gattaaagag cg 42
 43 <210> SEQ ID NO: 2
 44 <211> LENGTH: 36
 45 <212> TYPE: DNA
 46 <213> ORGANISM: Artificial Sequence
 48 <220> FEATURE:
 49 <223> OTHER INFORMATION: Description of Artificial Sequence: Flanking
 50 sequence of measles HA gene insert for N-P and P-M
 51 junctions.
 53 <400> SEQUENCE: 2
 54 gatccaacaa agaaacgaca ccgaacaaac cttaag 36
 57 <210> SEQ ID NO: 3
 58 <211> LENGTH: 101
 59 <212> TYPE: DNA
 60 <213> ORGANISM: Artificial Sequence
 62 <220> FEATURE:
 63 <223> OTHER INFORMATION: Description of Artificial Sequence: Flanking
 64 sequence of measles HA gene insert for HN-L
 65 junction.
 67 <400> SEQUENCE: 3
 68 aggcctaaaa gggaaatata aaaaacttag gagtaaagtt acgcaatcca actctactca 60

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74 <212> TYPE: DNA
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77 <220> FEATURE:
78 <223> OTHER INFORMATION: Description of Artificial Sequence: Flanking
79     sequence of measles HA gene insert for HN-L
80     junction.
82 <400> SEQUENCE: 4
83 tcataattaa ccataatatg catcaatota tctataatac aagtatatga taagtaatca 60
84 gcaatcagac aataggcct 79
87 <210> SEQ ID NO: 5
88 <211> LENGTH: 83
89 <212> TYPE: DNA
90 <213> ORGANISM: Artificial Sequence
92 <220> FEATURE:
93 <223> OTHER INFORMATION: Description of Artificial Sequence: Forward primer
94     for PCR of measles HA gene insert for N-P and P-M
95     junction.
97 <400> SEQUENCE: 5
98 ttaatcttaa gaatatacaa ataagaaaaa cttaggatta aagagcgatg tcaccacaac 60
99 gagaccggat aaatgccttc tac 83
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103 <211> LENGTH: 67
104 <212> TYPE: DNA
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108 <223> OTHER INFORMATION: Description of Artificial Sequence: Reverse primer
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110     junctions.
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113 attattgctt aaggtttggt cggtgtcggt tctttgttgg atcctatctg cgattggttc 60
114 catcttc 67
117 <210> SEQ ID NO: 7
118 <211> LENGTH: 55
119 <212> TYPE: DNA
120 <213> ORGANISM: Artificial Sequence
122 <220> FEATURE:
123 <223> OTHER INFORMATION: Description of Artificial Sequence: Forward primer
124     for PCR of measles HA gene insert for HN-L
125     junction.
127 <400> SEQUENCE: 7
128 gacaataggc ctaaaaggga aatataaaaa acttaggagt aaagttacgc aatcc 55
131 <210> SEQ ID NO: 8
132 <211> LENGTH: 68
133 <212> TYPE: DNA
134 <213> ORGANISM: Artificial Sequence
136 <220> FEATURE:

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137 <223> OTHER INFORMATION: Description of Artificial Sequence:
138     Reverse/Forward primer for PCR of measles HA gene
139     insert for HN-L junction.
141 <400> SEQUENCE: 8
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143 gtccttcc                                         68
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148 <212> TYPE: DNA
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152 <223> OTHER INFORMATION: Description of Artificial Sequence: Forward primer
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154     junction.
156 <400> SEQUENCE: 9
157 cagtcacccg ggaagatgga accaatcgca gatagtcata attaaccata atatgcatca 60
158 atctatctat aatacaa                                         77
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167 <223> OTHER INFORMATION: Description of Artificial Sequence: Reverse primer
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169     junction.
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176 <211> LENGTH: 28
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180 <220> FEATURE:
181 <223> OTHER INFORMATION: Description of Artificial Sequence:
182     Forward/Reverse primer for PCR of measles HA gene
183     insert for HN-L junction.
185 <400> SEQUENCE: 11
186 cggataaacg cggtctacaa agataacc                                         28
189 <210> SEQ ID NO: 12
190 <211> LENGTH: 25
191 <212> TYPE: DNA
192 <213> ORGANISM: Artificial Sequence
194 <220> FEATURE:
195 <223> OTHER INFORMATION: Description of Artificial Sequence: Reverse primer
196     for PCR of measles HA gene for N-P and M-P
197     junctions.
199 <400> SEQUENCE: 12
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203 <210> SEQ ID NO: 13
204 <211> LENGTH: 23

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205 <212> TYPE: DNA
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208 <220> FEATURE:
209 <223> OTHER INFORMATION: Description of Artificial Sequence: Upstream HPIV2
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212 <400> SEQUENCE: 13
213 gggccatgga agattacagc aat                                23
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217 <211> LENGTH: 25
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219 <213> ORGANISM: Artificial Sequence
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223     HPIV2 HN primer.
225 <400> SEQUENCE: 14
226 caataagctt aaagcattag ttccc                                25
229 <210> SEQ ID NO: 15
230 <211> LENGTH: 31
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232 <213> ORGANISM: Artificial Sequence
234 <220> FEATURE:
235 <223> OTHER INFORMATION: Description of Artificial Sequence: Upstream HPIV2
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238 <400> SEQUENCE: 15
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243 <211> LENGTH: 30
244 <212> TYPE: DNA
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247 <220> FEATURE:
248 <223> OTHER INFORMATION: Description of Artificial Sequence: Downstream
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251 <400> SEQUENCE: 16
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256 <211> LENGTH: 26
257 <212> TYPE: DNA
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260 <220> FEATURE:
261 <223> OTHER INFORMATION: Description of Artificial Sequence: HPIV1 HN
262     primer.
264 <400> SEQUENCE: 17
265 agtggctaata tgcattgcat ccacat                                26
268 <210> SEQ ID NO: 18
269 <211> LENGTH: 24
270 <212> TYPE: DNA
271 <213> ORGANISM: Artificial Sequence
273 <220> FEATURE:
274 <223> OTHER INFORMATION: Description of Artificial Sequence: HPIV1 HN

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277 <400> SEQUENCE: 18
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281 <210> SEQ ID NO: 19
282 <211> LENGTH: 15492
283 <212> TYPE: DNA
284 <213> ORGANISM: Artificial Sequence
286 <220> FEATURE:
287 <223> OTHER INFORMATION: Description of Artificial Sequence: Sequence of
288      pFLC.PIV32, 15492 bp in sense orientation.
290 <400> SEQUENCE: 19
291 accaaacaag agaagaaact tgtctgggaa tataaattta actttaaatt aacttaggat 60
292 taaagacatt gactagaagg tcaagaaaag ggaactctat aatttcaaaa atgttgagcc 120
293 tatttgatac atttaatgca cgtaggcaag aaaacataac aaaatcagcc ggtggagcta 180
294 tcattcctgg acagaaaaat actgtctcta tattcgccct tggaccgaca ataactgatg 240
295 ataatgagaa aatgacatta gctcttctat ttctatctca ttactagat aatgagaaac 300
296 aacatgcaca aagggcaggg ttcttgggtg ttttattgtc aatggcttat gccaatccag 360
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323 aaacgacaca atcaacacaa gaaccagca actcagtgcc accatctgtc aaccagaaat 1980
324 caaaccaaca gaaacaagt agaaagatag tggatcaact gacaaaaata gacagtccgg 2040
325 gtcatcacac gaatgtacaa cagaagcaaa agatagaaat attgatcagg aaactgtaca 2100
326 gagaggacct gggagaagaa gcagctcaga tagtagagct gagactgtgg tctctggagg 2160
327 aatccccaga agcatcacag attctaaaaa tggaaccaa aacacggagg atattgatct 2220
328 caatgaaatt agaaagatgg ataaggactc tattgagggg aaaatgcgac aatctgcaaa 2280

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